RAW SEQUENCE LISTING PATENT APPLICATION US/08/734,443

DATE: 01/27/98 TIME: 13:50:28

INPUT SET: S22894.raw

This Raw Listing contains the General Information Section and up to the first 5 pages.

1	SEQUENCE LISTING		
2			
3	(1) Ger	eral Information:	
4 5 6	(i) <i>P</i>	APPLICANT: Keyt, Bruce A. Nguyen, Francis H. Ferrara, Napoleone	
7 8		reflata, Napoleone	
9 10	(ii) 7	TITLE OF INVENTION: Variants of Vascular Endothelial Cell Growth Factor Having Antagonistic Properties	
11			
12	(iii) N	JUMBER OF SEQUENCES: 14	
13 14	(iv) (CORRESPONDENCE ADDRESS: (A) ADDRESSEE: Walter H. Dreger	
15		(B) STREET: 4 Embarcadero Center, Suite 3400	
16 17		(C) CITY: San Francisco	
18		(D) STATE: California	
19		(E) COUNTRY: United States	
20		(F) ZIP: 94111	
21			
22	(V) (COMPUTER READABLE FORM:	
23		(A) MEDIUM TYPE: Floppy disk	
24		(B) COMPUTER: IBM PC compatible	
25		(C) OPERATING SYSTEM: PC-DOS/MS-DOS	
26		(D) SOFTWARE: PatentIn Release #1.0, Version #1.30	
27			
28	(Vi) (CURRENT APPLICATION DATA:	
29		(A) APPLICATION NUMBER: US 08/734,443	
30		(B) FILING DATE: 17-OCT-1996	
31		(C) CLASSIFICATION:	
32	41111	AMMODNEY/AGENM INFORMATION.	
33	(V111) .	ATTORNEY/AGENT INFORMATION: (A) NAME: Dreger, Walter H.	
34		(B) REGISTRATION NUMBER: 24,190	
35 36		(C) REFERENCE/DOCKET NUMBER: A-63096/WHD/MTK	
37		(C) KEFERENCE, BOCKET ROLLDER II OCCUPATION	
38	(iv)	TELECOMMUNICATION INFORMATION:	
39	(14)	(A) TELEPHONE: 415-781-1989	
40		(B) TELEFAX: 415-398-3249	
41			
42			
43	(2) INFOR	MATION FOR SEQ ID NO:1:	
44			
45	(i)	SEQUENCE CHARACTERISTICS:	
46		(A) LENGTH: 59 base pairs	

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47 48 49 50	(B) TYPE: nucleic acid(C) STRANDEDNESS: single(D) TOPOLOGY: linear	
51 52 53 54 55	<pre>(ii) MOLECULE TYPE: other nucleic acid (A) DESCRIPTION: /desc = "oligonucleotide"</pre>	
56 57 58	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:1:	
59 60 61 62	CCTATGGCTG AAGGCGGCCA GAAGCCTCAC GAAGTGGTGA AGTTCATGGA CGTGTATCA	59
63 64	(2) INFORMATION FOR SEQ ID NO:2:	
65 66 67 68 69	 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 30 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 	
71 72 73	<pre>(ii) MOLECULE TYPE: other nucleic acid (A) DESCRIPTION: /desc = "oligonucleotide"</pre>	
7 4 75	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:2:	
76 77 78 79	AGTAGCAAGC TTGACGTGTG GCAGGCTTGA	30
80 81	(2) INFORMATION FOR SEQ ID NO:3:	
82 83 84 85 86 87	 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 29 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 	
88 89 90	<pre>(ii) MOLECULE TYPE: other nucleic acid (A) DESCRIPTION: /desc = "oligonucleotide"</pre>	
91 92	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:3:	
93 94 95 96	GATCTGGCCA TACACTTGAG TGACAATGA	29
97 98	(2) INFORMATION FOR SEQ ID NO:4:	
99	(i) SEQUENCE CHARACTERISTICS:	

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	INPUT SET: S22894.rav
(A) LENGTH: 28 base pairs	
(B) TYPE: nucleic acid	
(C) STRANDEDNESS: single	
(D) TOPOLOGY: linear	
(ii) MOLECULE TYPE: other nucleic acid	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:4:	
-	
CATCCACTTT GCCTTTCTCT CCACAGGT	28
(2) INFORMATION FOR SEC ID NO:5:	
(2) INFORMATION FOR DEG 1D NO.3.	
(1) CECHENCE CUNDACTEDICTICS.	
, , =	
· ·	
(D) TOPOLOGY: linear	
(A) DESCRIPTION: /desc = "oligonucleotide"	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:5:	
GTCCACTCCC AG	12
(2) INFORMATION FOR SEQ ID NO:6:	
(i) SEQUENCE CHARACTERISTICS:	
(A) LENGTH: 30 base pairs	
(B) TYPE: nucleic acid	
(C) STRANDEDNESS: single	
(D) TOPOLOGY: linear	
(ii) MOLECULE TYPE: other nucleic acid	
(A) DESCRIPTION: /desc = "oligonucleotide"	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:6:	
• • •	
AGGCTGCTGC AGTTCGACGT GGGAGTGGAC	30
(2) INFORMATION FOR SEO ID NO:7:	
(a) Torrantation torrang to motive	
(i) SECTIENCE CHARACTERISTICS.	
• • •	
(D) TIPE: NUCLEIC ACID	
	(B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear (ii) MOLECULE TYPE: other nucleic acid (A) DESCRIPTION: /desc = "oligonucleotide" (xi) SEQUENCE DESCRIPTION: SEQ ID NO:4: CATCCACTTT GCCTTTCTCT CCACAGGT (2) INFORMATION FOR SEQ ID NO:5: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 12 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear (ii) MOLECULE TYPE: other nucleic acid (A) DESCRIPTION: /desc = "oligonucleotide" (xi) SEQUENCE DESCRIPTION: SEQ ID NO:5: GTCCACTCCC AG (2) INFORMATION FOR SEQ ID NO:6: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 30 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear (ii) MOLECULE TYPE: other nucleic acid (A) DESCRIPTION: /desc = "oligonucleotide" (xi) SEQUENCE DESCRIPTION: /desc = "oligonucleotide" (xi) SEQUENCE DESCRIPTION: SEQ ID NO:6:

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		1NFUL SEL: 344074.748
153 154	(C) STRANDEDNESS: single(D) TOPOLOGY: linear	
155		
156	(ii) MOLECULE TYPE: other nucleic acid	
157	<pre>(A) DESCRIPTION: /desc = "oligonucleotide"</pre>	
158		
159	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:7:	
160		
161	CAGGGGCACA TCGGATGGCT TGAA	24
162		
163		
164		
165	(2) INFORMATION FOR SEQ ID NO:8:	
166		
167	(i) SEQUENCE CHARACTERISTICS:	
168	(A) LENGTH: 24 base pairs	
169	(B) TYPE: nucleic acid	
170	(C) STRANDEDNESS: single	
171	(D) TOPOLOGY: linear	
172		
173	(ii) MOLECULE TYPE: other nucleic acid	
174	<pre>(A) DESCRIPTION: /desc = "oligonucleotide"</pre>	
175		
176	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:8:	
177		
178	CAGGGGCACG GCGGATGGCT TGAA	24
179		
180		
181		
182		
183		
184		
185	(2) INFORMATION FOR SEQ ID NO:9:	
186		
187	(i) SEQUENCE CHARACTERISTICS:	
188	(A) LENGTH: 24 base pairs	
189	(B) TYPE: nucleic acid	
190	(C) STRANDEDNESS: single	
191	(D) TOPOLOGY: linear	
192	(ii) vornourn munn sellere med de seid	
193	(ii) MOLECULE TYPE: other nucleic acid	
194	(A) DESCRIPTION: /desc = "oligonucleotide"	
195	/ GEOMENGE DEGGDIDETON: GEO. ID NO. 0.	
196	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:9:	
197	CHICAPHICOAA HOOOOGOOO AHOO	24
198	GTCATTGCAA TCGCCCCCGC ATCG	24
199		
200		
201 202	(2) INFORMATION FOR CEO IN MO. 10.	
202	(2) INFORMATION FOR SEQ ID NO:10:	
203	(i) SEQUENCE CHARACTERISTICS:	
204	(A) LENGTH: 24 base pairs	
200	(v) prugin. 54 page barra	

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		INPUT SET: S22	894.ra
206	(B) TYPE: nucleic acid		
207	(C) STRANDEDNESS: single		
208	(D) TOPOLOGY: linear		
209			
210	(ii) MOLECULE TYPE: other nucleic acid		
211	(A) DESCRIPTION: /desc = "oligonucleotide"		
212			
213	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:10:		
214			
215	GTCATTGCAG GCGCCCCGC ATCG		24
216			
217			
218	10. 700000000000000000000000000000000000		
219	(2) INFORMATION FOR SEQ ID NO:11:		
220	() CHOUSING COURT COURT		
221	(i) SEQUENCE CHARACTERISTICS:		
222	(A) LENGTH: 51 base pairs		
223	(B) TYPE: nucleic acid		
224	(C) STRANDEDNESS: single		
225 226	(D) TOPOLOGY: linear		
226	(ii) NOIEGUE EUDE, other musleis esid		
227	(ii) MOLECULE TYPE: other nucleic acid		
228	(A) DESCRIPTION: /desc = "oligonucleotide"		
230	(vi) CECULENCE DECORPORATION, CEC ID NO.11.		
231	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:11:		
232	GTCATTGCAG GCGCCCCCGC ATCGCATCAG GGGCACGGCG GATGGCTTG	n n	51
233	GICHITGERG GEGECECEGE ATEGERTERG GGGCREGGEG GRIGGETTO	ч н	31
234			
235			
236	(2) INFORMATION FOR SEQ ID NO:12:		
237	(-,		
238	(i) SEQUENCE CHARACTERISTICS:		
239	(A) LENGTH: 51 base pairs		
240	(B) TYPE: nucleic acid		
241	(C) STRANDEDNESS: single		
242	(D) TOPOLOGY: linear		
243	, <i>,</i>		
244	(ii) MOLECULE TYPE: other nucleic acid		
245	(A) DESCRIPTION: /desc = "oligonucleotide"		
246	<u>-</u>		
247	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:12:		
248			
249	GTCATTGCAA TCGCCCCCGC ATCGCATCAG GGGCACATCG GATGGCTTG	A A	51
250			
251			
252			
253	(2) INFORMATION FOR SEQ ID NO:13:		
254			
255	(i) SEQUENCE CHARACTERISTICS:		
256	(A) LENGTH: 990 base pairs		
257	(B) TYPE: nucleic acid		
258	(C) STRANDEDNESS: double		

SEQUENCE VERIFICATION REPORT PATENT APPLICATION *US/08/734,443*

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Original Text

379

Stop Codon at end of sequence removed - no error